Figure 1

1	GCCTCTCCCATCAGACACCCCAAGGTTCC								
Τ.	CGGAGAGAGGTAGTCTGTGGGGTTCCAAGG								
31	ATCCGAAGCAGGCGGAGCACCGAACGCACC								
31	TAGGCTTCGTCCGCCTCGTGGCTTGCGTGG	60							
61	CCGGGTGGTCAGGGACCCCCATCCGTGCT	90							
O 1.	GGCCCCACCAGTCCCTGGGGGTAGGCACGA	90							
91	GCCCCTAGGAGCCCGCGCCTCTCCTGC	120							
91	CGGGGGATCCTCGGGCGCGGAGAGAGACG	120							
121	GCCCGCCTCTCGGGCCGCAACATCGCGCG	150							
<u> </u>	CGGGGCGAGAGCCCGGCGTTGTAGCGCGC	100							
151	GTTCCTTTAACAGCGCGCTGGCAGGGTGTG	180							
T	CAAGGAAATTGTCGCGCGACCGTCCCACAC	100							
181	GGAAGCAGGACCGCGTCCTCCCGCCCCCTC	210							
ТОТ	CCTTCGTCCTGGCGCAGGAGGCGGGGGAG	Z .L (
211	CCATCCGAGTTTCAGGTGAATTGGTCACCG	240							
Z 1 1	GGTAGGCTGAAAGTCCACTTAACCAGTGGC	470							

0.4.7	AGGGAGGAGCCGACACCACACCTACAC										
241	TCCCTCCTCCGGCTGTGTGGTGTGGATGTG	271									
	TCCCGCGTCCACCTCTCCCTGCTTCC	300									
271	AGGGCGCAGGTGGAGAGGGACGAAGG	300									
301	TCTTGGCGGAGGCGCAGGAACCGAGAGCC	330									
201	AGAACCGCCTCCGCCGTCCTTGGCTCTCGG	330									
331	AGGTCCAGAGCGCCGAGGAGCCGGTCTAGG	360									
221	TCCAGGTCTCGCGGCTCCTCGGCCAGATCC	300									
361	ACGCAGCAGATTGGTTTATCTTGGAAGCTA	390									
201	TGCGTCGTCTAACCAAATAGAACCTTCGAT	330									
391	AAGGCATTGCTCATCCTGAAGATCAGCTG	420									
J J I	TTCCCGTAACGAGTAGGACTTCTAGTCGAC START	120									
421	ACCATTGACAATCAGCCATGTCATCCAGGC	450									
	TGGTAACTGTTAGTCGGTACAGTAGGTCCG M S S R P										
	CTCTTGAAAGTCCACCTCCTTACAGGCCTG	480									
451	GAGAACTTTCAGGTGGAGGAATGTCCGGAC L E S P P P Y R P D	70U									

481	ATGAATTCAAACCGAATCATTATGCACCAA									
	TACTTAAGTTTGGCTTAGTAATACGTGGTT E F K P N H Y A P S	510								
511	GCAATGACATATATGGTGGAGAGATGCATG	540								
541	TTCGACCAATGCTCTCTCAGCCAGCCTACT AAGCTGGTTACGAGAGAGTCGGTCGGATGA R P M L S Q P A Y S	570								
571	CTTTTTACCCAGAAGATGAAATTCTTCACT GAAAAATGGGTGTTCTACTTTAAGAAGTGA F Y P E D E I L H F	600								
601	TCTACAAATGGACCTCTCCTCCAGGAGTGA AGATGTTTACCTGGAGAGGAGGTCCTCACT Y K W T S P P G V I	630								
631	TTCGGATCCTGTCTATGCTCATTATTGTGA									
661	TGTGCATTGCCATCTTTGCCTGTGTGGCC	690								

CC	AC(GCT'	rgc(CTG	GGA(CAG	AGG(CTA	rgg <i>i</i>	A.A	7
GG	TG	 CGA	 ACG	 GAC	CCT	GTC	TCC	GAT	 ACC'	$^{}$	/
	Τ	L	A	W	D	R	G	Y	G	T	
CI	TC	ССТ 	TTT 	AGG	AGG	TAG	TGT	AGG 	CTA	CC	-
GP	AG	GGA	AAA	TCC	TCC	ATC	ACA	TCC	GAT	GG	,
	S	L	L	G	G	S	V	G	Y	P	
CI	TA	TGG 	AGG 	AAG 	TGG	CTT 	TGG 	TAG 	CTA 	CG	-
GA	AT.	ACC	TCC	TTC	ACC	GAA	ACC	ATC	GAT	GC	,
	Y	G	G	S	G	F	G	S	Y	G	
GA	\AG	TGG 	CTA 	.TGG 	CTA 	TGG 	CTA 	TGG 	TTA 	TG 	{
СЛ	TC	ACC	GAT	ACC	GAT	ACC	GAT	ACC	AAT.	AC	
	S	G	Y	G	Y	G	Y	G	Y	G	
G(CTA 	TGG	CTA 	.CGG 	AGG	CTA	TAC	AGA 	.CCC.	AA 	{
CC	TAE	ACC	GAT	GCC	TCC	GAT	ATG	TCT	GGG	TT	•
	Y	G	Y	G	G	Y	Т	D	Р	R	
GF	4GC	AGC	AAA	.GGG	CTT	'CAT	'GTT	GGC	CAT	GG	,
CI	 ГСG	 TCG		 CCC'	 'GAA	 GTA.	 .CAA	 CCG	 GTA	 CC	8
	A	A	K	G	F	M	L	A	M	À	
C	rgc	CTT	'TTG	TTT	'CAT	'TGC	:CGC	GTT	'GGT	GA	
	 1 (G	 Gad	 \AAC	 \a\a\a	 ata	 ACG	 :GCG	 AAD.	 .CCA	 Ст	-
<u></u>					-				V	_	

901	TCTTTGTTACCAGTGTTATAAGATCTGAAA										
	AGAAACAATGGTCACAATATTCTAGACTTT F V T S V I R S E M	930									
931	TGTCCAGAACAAGAAGATACTACTTAAGTG	960									
	ACAGGTCTTGTTCTTCTATGATGAATTCAC S R T R R Y Y L S V										
961	TGATAATAGTGAGTGCTATCCTGGGCATCA	990									
	ACTATTATCACTCACGATAGGACCCGTAGT I I V S A I L G I M										
991	TGGTGTTTATTGCCACAATTGTCTATATAA	1020									
	ACCACAAATAACGGTCTTAACAGATATATT V F I A T I V Y I M										
1021	TGGGAGTGAACCCAACTGCTCAGTCTTCTG	1050									
	ACCCTCACTTGGGTTGACGAGTCAGAAGAC G V N P T A Q S S G										
1051		1080									
	CTAGAGATATACCAAGTCTTTATATACGGG S L Y G S Q I Y A L										
1081		1110									
	AGACGTTGGTTAAAATATGTGGACGTCGAT C N Q F Y T P A A T										

1111	CTGGACTCTACGTGGATCAGTATTTGTATC								
	GACCTGAGATCGACCTAGTCATAAACATAG G L Y V D Q Y L Y H								
1141	ACTACTGTGTTGTGGATCCCCAGGAGGCCA TGATGACACAACACCTAGGGGTCCTCCGGT Y C V D P Q E A I	1170							
1171	TTGCCATTGTACTGGGGTTCATGATTATTG AACGGTAACATGACCCCAAGTACTAATAAC A I V L G F M I I V	1200							
1201	TGGCTTTTGCTTTAATAATTTTCTTTGCTG ACCGAAAACGAAATTATTAAAAGAAACGAC A F A L I I F F A V	1230							
1231	TGAAAACTCGAAGAAAGATGGACAGGTATG ACTTTTGAGCTTCTTTCTACCTGTCCATAC K T R R K M D R Y D	1260							
1261	ACAAGTCCAATATTTTGTGGGACAAGGAAC TGTTCAGGTTATAAAACACCCTGTTCCTTG K S N I L W D K E H	1290							
1291	ACATTTATGATGAGCAGCCCCCAATGTCG TGTAAATACTACTCGTCGGGGGGTTACAGC	1320							

1 2 0 1	AGGAGTGGGTTAAAAATGTGTCTGCAGGCA													
1321	TCCT	CAC	CCA	'TTA	TTT	ACA	CAG.	ACG	TCC(G		1350			
1351	CACA	GGA(TTC. 				TGA(1380			
	GTGT ^e Q								ACT(D					
1381	ATGT										1410			
	ATCA V								CCG' A					
1411	ACTC	TTC			CAA 				CAA	GC	1440			
T.4.T.T.	TGAG.		GTT.	ACC	GTT	TCA	CTT	ACT	GTT K		1440			
1441	GGTT	TTA' 	TCC.	AGA 	GTC 	TTC 	CTA 	TAA 	ATC	CA 	1470			
	CCAA F								TAG S		11,0			
1471	CGCC	GGT	TCC 	TGA 	AGT	GGT	TCA	.GGA			1500			
	GCGG P									AG				
1501	CATT	'AAC 	TTC 	GCC	TGT 	'GGA 	.TGA 	.CTT	'CAG		1530			
- 	GTAA L								GTC R					

1531	AGCCTCGTTACAGCAGCGGTGGTAACTTTG									
1231	TCGGAGCAATGTCGTCGCCACCATTGAAAC PRYSSGGNFE	1560								
1561	AGACACCTTCAAAAAGAGCACCTGCAAAGG TCTGTGGAAGTTTTCTCGTGGACGTTTCC T P S K R A P A K G	1590								
1591	GAAGAGCAGGAAGGTCAAAGAGAACAGAGC CTTCTCGTCCTTCCAGTTTCTCTTGTCTCG R A G R S K R T E Q	1620 Q								
1621	AAGATCACTATGAGACAGACTACACAACTG TTCTAGTGATACTCTGTCTGATGTGTTGAC D H Y E T D Y T T G	1650								
1651	GCGGCGAGTCCTGTGATGAGCTGGAGGAGG CGCCGCTCAGGACACTACTCGACCTCCTCC G E S C D E L E E D	1680								
1681	ACTGGATCAGGGAATATCCACCTATCACTT TCACCTAGTCCCTTATAGGTGGATAGTGAA W I R E Y P P I T S	1710								
1711	CAGATCAACAAAGACAACTGTACAAGAGGAGTCTAGTTGTTTCTCTTTGACATGTTCTCCT	1740								

7 7 4 7	ATTTTGACACTGGCCTACAGGAATACAAGA										
1741	TAAAAC		ACCG	GATG	TCCT	TATO			1770		
1	GCTTAC	CAATC	AGAA	.CTTG	ATGA	GAT(CAAT	A	1000		
1771	CGAATO	GTTAG' S							1800		
1801	AAGAAC	TCTC(1830		
	TTCTTC E I	GAGAG L S	-								
1831	ATGACT	[ATAG	AGAA 	.GAA.	GTGA	AGA(GTAC 	!A -	1860		
	TACTGA D	ATATC Y R									
1861	TGGCT				. – – –			-	1890		
	ACCGA(CGACG A A									
1891	AGCAA(GCAGA				1920		
	TCGTT(CACTT V K									
1921	GTAAG				AAGCA 				1950		
	CATTC'	TTCTT K N									

1951	GCAAATTGTCACACATCAAGAAGATGGTTG	1000
1951	CGTTTAACAGTGTGTAGTTCTTCTACCAAC K L S H I K K M V G STOP	1980
1981	GAGACTATGATAGACAGAAAACATAGAAGG	2010
2011	D Y D R Q K T CTGATGCCAAGTTGTTTGAGAAATTAAGTA GACTACGGTTCAACAAACTCTTTAATTCAT	2040
2041	TCTGACATCTCTGCAATCTTCTCAGAAGGC AGACTGTAGAGACGTTAGAAGAGTCTTCCG	2070
2071	AAATGACTTTGGACCATAACCCCGGAAGCC TTTACTGAAACCTGGTATTGGGGCCTTCGG	2100
2101	AAACCTCTGTGAGCATCACAAAGTTTTGGGGGGAGACACTCGTAGTGTTTCAAAACCC	3 2130
2131	TTGCTTTAACATCATCAGTATTGAAGCATT AACGAAATTGTAGTAGTCATAACTTCGTAA	2160
2161	TTATAAATCGCTTTTGATAATCAACTGGGCAATATTTAGCGAAAACTATTAGTTGACCCG	2190

11/14

2191	TGAACAACTCCAATTAAGGATTTTATGCTT						
	ACTTGTTGAGGTTAATTCCTAAAATACGAA	2220					
2221	TAAACATTGGTTCTTGTATTAAGAATGAAA	2250					
	ATTTGTAACCAAGAACATAATTCTTACTTT						
2251	TACTGTTTGAGGTTTTTTAAGCCTTAAAGGA	2280					
	ATGACAAACTCCAAAAATTCGGAATTTCCT						
2281	AGGTTCTGGTGTGAACTAAACTTTCACACC	2310					
	TCCAAGACCACATTGATTTGAAAGTGTGG						
2311	CC 	2312					
	GG						

Figure 2

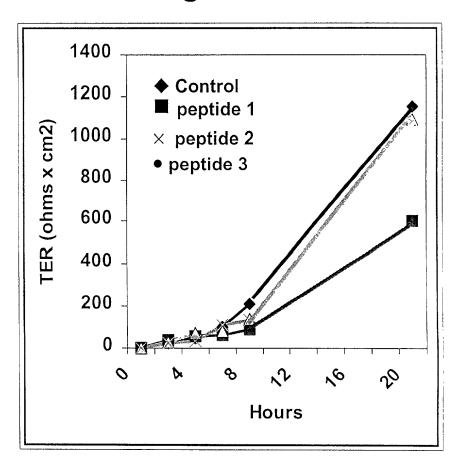


Figure 3A

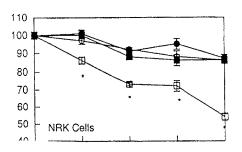


Figure 3B

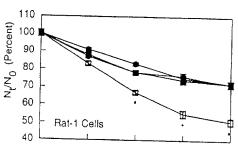
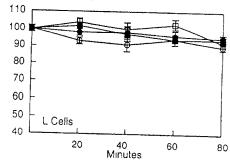


Figure 3C



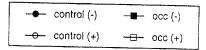


Figure 4

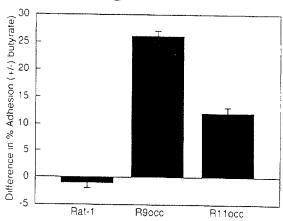


Figure 5A

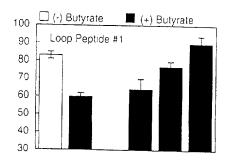


Figure 5B

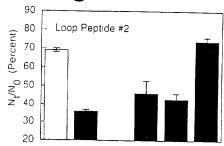


Figure 5C

